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OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/730,374

DATE: 05/24/2001
TIME: 15:13:14

Input Set : A:\150188us2.txt
Output Set: C:\CRF3\05242001\I730374.raw

ENTERED

5 <110> APPLICANT: Lust, John A.
 6 Donovan, Kathleen A.
 8 <120> TITLE OF INVENTION: USE OF GENETICALLY ENGINEERED ANTIBODIES
 9 TO CD38 TO TREAT MULTIPLE MYELOMA
 12 <130> FILE REFERENCE: 150.188US2
 14 <140> CURRENT APPLICATION NUMBER: 09/730,374
 C--> 15 <141> CURRENT FILING DATE: 2001-05-10
 17 <150> PRIOR APPLICATION NUMBER: PCT/US99/12512
 18 <151> PRIOR FILING DATE: 1999-06-04
 20 <150> PRIOR APPLICATION NUMBER: 60/088,277
 21 <151> PRIOR FILING DATE: 1998-08-05
 23 <160> NUMBER OF SEQ ID NOS: 4
 25 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 27 <210> SEQ ID NO: 1
 28 <211> LENGTH: 750
 29 <212> TYPE: DNA
 30 <213> ORGANISM: Artificial Sequence
 32 <220> FEATURE:
 33 <223> OTHER INFORMATION: A nucleotide sequence encoding a single chain
 34 variable region fragment (scFv)
 36 <400> SEQUENCE: 1
 37 ggcgcggccg gccatggcca aggtccagct gcaggaggta ggaccttagcc tagtgcagcc 60
 38 ctcacagcgc ctgtccataa cctgcacagt ctctggttc tcattaattt gttatggtgt 120
 39 acactgggtt cgccagtctc caggaaaggg tctggagtgg ctgggagtgat tatggagagg 180
 40 tggaaagcaca gactacaatg cagcttcat gtccagactg agcatcacca aggacaactc 240
 41 caagagccaa gttttcttta aaatgaacag tctgcaagct gatgacactg ccatataactt 300
 42 ctgtgccaaa accttgatta cgacgggcta tgctatggac tactggggcc aagggaccac 360
 43 ggtcaccgtc tcctcagggtg gaggcggttc aggccggaggt ggctctggcg gtggcggatc 420
 44 ggacatcgag ctcactcagt ctccatcctc cttttctgtt tctcttaggag acagagtcac 480
 45 cattacttgc aaggcaagtg aggacatata taatcggtt gcctggatc agcaagaaacc 540
 46 aggaaatgtt cctaggctct taatatctgg tgcaaccagt ttggaaactg gggttccttc 600
 47 aagattcagt ggcagtggat ctggaaaggta ttacactctc agcattacca gtcttcagac 660
 48 tgaagatgtt gctacttatt actgtcaaca gtattggagt actcctacgt tcggtgagg 720
 49 gaccaagctg gaaatcaaacc gggcgccgc 750
 51 <210> SEQ ID NO: 2
 52 <211> LENGTH: 241
 53 <212> TYPE: PRT
 54 <213> ORGANISM: Artificial Sequence
 56 <220> FEATURE:
 57 <223> OTHER INFORMATION: A polypeptide encoded by an open reading frame of
 58 SEQ ID NO:1
 60 <400> SEQUENCE: 2
 61 Gly Pro Ala Gly His Gly Gln Gly Pro Ala Ala Gly Val Arg Thr Pro
 62 1 5 10 15
 63 Ser Ala Ala Leu Thr Ala Pro Val His Asn Leu His Ser Leu Trp Phe
 64 20 25 30
 65 Leu Ile Asn Leu Trp Cys Thr Leu Gly Ser Pro Val Ser Arg Lys Gly

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66      35          40          45
67 Ser Gly Val Ala Gly Ser Asp Met Glu Arg Trp Lys His Arg Leu Gln
68      50          55          60
69 Cys Ser Phe His Val Gln Thr Glu His His Gln Gly Gln Leu Gln Glu
70      65          70          75          80
71 Pro Ser Phe Leu Asn Glu Gln Ser Ala Ser His Cys His Ile Leu Leu
72      85          90          95
73 Cys Gln Asn Leu Asp Tyr Asp Gly Leu Cys Tyr Gly Leu Leu Gly Pro
74      100         105         110
75 Arg Asp His Gly His Arg Leu Leu Arg Trp Arg Arg Phe Arg Arg Arg
76      115         120         125
77 Trp Leu Trp Arg Trp Arg Ile Gly His Arg Ala His Ser Val Ser Ile
78      130         135         140
79 Leu Leu Phe Cys Ile Ser Arg Arg Gln Ser His His Tyr Leu Gln Gly
80      145         150         155         160
81 Lys Gly His Ile Ser Val Ser Leu Val Ser Ala Glu Thr Arg Lys Cys
82      165         170         175
83 Ser Ala Leu Asn Ile Trp Cys Asn Gln Phe Gly Asn Trp Gly Ser Phe
84      180         185         190
85 Lys Ile Gln Trp Gln Trp Ile Trp Lys Gly Leu His Ser Gln His Tyr
86      195         200         205
87 Gln Ser Ser Asp Arg Cys Cys Tyr Leu Leu Leu Ser Thr Val Leu Glu
88      210         215         220
89 Tyr Ser Tyr Val Arg Trp Arg Asp Gln Ala Gly Asn Gln Thr Gly Gly
90      225         230         235         240
91 Arg
94 <210> SEQ ID NO: 3
95 <211> LENGTH: 249
96 <212> TYPE: PRT
97 <213> ORGANISM: Artificial Sequence
99 <220> FEATURE:
100 <223> OTHER INFORMATION: A polypeptide encoded by an open reading frame of
101     SEQ ID NO:1
104 <400> SEQUENCE: 3
105 Ala Gln Pro Ala Met Ala Lys Val Gln Leu Gln Glu Ser Gly Pro Ser
106     1           5           10          15
107 Leu Val Gln Pro Ser Gln Arg Leu Ser Ile Thr Cys Thr Val Ser Gly
108     20          25          30
109 Phe Ser Leu Ile Ser Tyr Gly Val His Trp Val Arg Gln Ser Pro Gly
110     35          40          45
111 Lys Gly Leu Glu Trp Leu Gly Val Ile Trp Arg Gly Gly Ser Thr Asp
112     50          55          60
113 Tyr Asn Ala Ala Phe Met Ser Arg Leu Ser Ile Thr Lys Asp Asn Ser
114     65          70          75          80
115 Lys Ser Gln Val Phe Phe Lys Met Asn Ser Leu Gln Ala Asp Asp Thr
116     85          90          95
117 Ala Ile Tyr Phe Cys Ala Lys Thr Leu Ile Thr Thr Gly Tyr Ala Met
118     100         105         110
119 Asp Tyr Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Gly Gly Gly

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120      115          120          125
121 Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser Asp Ile Glu Leu
122      130          135          140
123 Thr Gln Ser Pro Ser Ser Phe Ser Val Ser Leu Gly Asp Arg Val Thr
124      145          150          155          160
125 Ile Thr Cys Lys Ala Ser Glu Asp Ile Tyr Asn Arg Leu Ala Trp Tyr
126      165          170          175
127 Gln Gln Lys Pro Gly Asn Ala Pro Arg Leu Leu Ile Ser Gly Ala Thr
128      180          185          190
129 Ser Leu Glu Thr Gly Val Pro Ser Arg Phe Ser Gly Ser Gly
130      195          200          205
131 Lys Asp Tyr Thr Leu Ser Ile Thr Ser Leu Gln Thr Glu Asp Val Ala
132      210          215          220
133 Thr Tyr Tyr Cys Gln Gln Tyr Trp Ser Thr Pro Thr Phe Gly Gly Gly
134      225          230          235          240
135 Thr Lys Leu Glu Ile Lys Arg Ala Ala
136      245
138 <210> SEQ ID NO: 4
139 <211> LENGTH: 239
140 <212> TYPE: PRT
141 <213> ORGANISM: Artificial Sequence
143 <220> FEATURE:
144 <223> OTHER INFORMATION: A polypeptide encoded by an open reading frame of
145     SEQ ID NO:1
148 <400> SEQUENCE: 4
149 Pro Ser Arg Pro Trp Pro Arg Ser Ser Cys Arg Ser Gln Asp Leu Ala
150 1          5          10          15
151 Cys Ser Pro His Ser Ala Cys Pro Pro Ala Gln Ser Leu Val Ser His
152 20         25         30
153 Leu Val Met Val Tyr Thr Gly Phe Ala Ser Leu Gln Glu Arg Val Trp
154 35         40         45
155 Ser Gly Trp Glu Tyr Gly Glu Val Glu Ala Gln Thr Thr Met Gln Leu
156 50         55         60
157 Ser Cys Pro Asp Ala Ser Pro Arg Thr Thr Pro Arg Ala Lys Phe Ser
158 65         70         75         80
159 Leu Lys Thr Val Cys Lys Leu Met Thr Leu Pro Tyr Thr Ser Val Pro
160 85         90         95
161 Lys Pro Leu Arg Arg Ala Met Leu Trp Thr Thr Gly Ala Lys Gly Pro
162 100        105        110
163 Arg Ser Pro Ser Pro Gln Val Glu Ala Val Gln Ala Glu Val Ala Leu
164 115        120        125
165 Ala Val Ala Asp Arg Thr Ser Ser Ser Leu Ser Leu His Pro Pro Phe
166 130        135        140
167 Leu Tyr Leu Glu Thr Glu Ser Pro Leu Leu Ala Arg Gln Val Arg Thr
168 145        150        155        160
169 Tyr Ile Ile Gly Pro Gly Ile Ser Arg Asn Gln Glu Met Leu Leu Gly
170 165        170        175
171 Ser Tyr Leu Val Gln Pro Val Trp Lys Leu Gly Phe Leu Gln Asp Ser
172 180        185        190

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173 Val Ala Val Asp Leu Glu Arg Ile Thr Leu Ser Ala Leu Pro Val Phe
174 195 200 205
175 Arg Leu Lys Met Leu Leu Ile Thr Val Asn Ser Ile Gly Val Leu
176 210 215 220
177 Leu Arg Ser Val Glu Gly Pro Ser Trp Lys Ser Asn Gly Arg Pro
178 225 230 235

VERIFICATION SUMMARY
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L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date